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Key indicators

Single-crystal X-ray study

T = 295 K

Mean $\sigma(\text{C}-\text{C}) = 0.007 \text{ \AA}$

R factor = 0.040

wR factor = 0.101

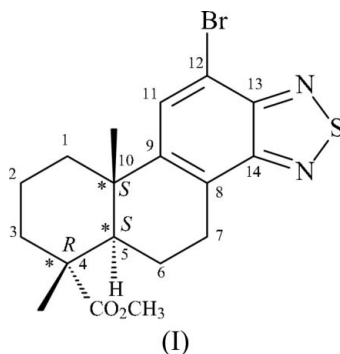
Data-to-parameter ratio = 13.9

For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Methyl 12-bromo-13,14-(thiodinitrilo)-
deisopropyldehydroabietateThe structure of the title compound, $\text{C}_{18}\text{H}_{21}\text{BrN}_2\text{O}_2\text{S}$, features
a five-membered 2,1,3-thiadiazole ring in which the N atoms
are two-coordinate [$\text{N}-\text{S}-\text{N} = 101.6 (2)^\circ$].

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Comment

The reaction of methyl 12-bromo-13,14-dinitrodeisopropyl-
dehydroabietate, the structure of which was reported recently
(Pan *et al.*, 2005), with thionyl chloride leads to the formation
of methyl 12-bromo-13,14-(2,1,3)-thiadiazolydehydroabietate,
(I) (Fig. 1).The bond dimensions of the five-membered ring in (I)
compare well with those found in related compounds, such as
benzo-2,1,3-thiazole (Luzzati, 1951) and 3,4-diphenyl-1,2,5-
thiadiazole (Mellini & Merlino, 1976). The bond dimensions in
the other part of the molecule are similar to those of 12-
bromo-13,14-dinitrodeisopropyldehydroabietate (Pan *et al.*,
2005) itself.

Experimental

12-Bromo-13,14-diaminodeisopropyldehydroabietate (Fonseca *et al.*,
2004) (1.01 g, 2.65 mmol) was dissolved in dry benzene (15 ml) and
the solution was cooled to 273 K. To the solution was added thionyl
chloride (0.81 g, 6.84 mmol) dissolved in benzene (15 ml). The
mixture was stirred for 0.5 h at 273 K, and then at room temperature
for another 1 h before being further refluxed for 12 h. The solvent
was removed and the residue was purified by column chromatography
on silica gel (light petroleum–ethyl acetate 5:1 *v/v*) to give
pale-yellow crystals of (I) (0.423 g, 1.034 mmol, 40% yield; mp 370–
371 K). Single crystals of (I) were obtained from an absolute acetone
solution.

Crystal data

$C_{18}H_{21}BrN_2O_2S$
 $M_r = 409.34$
 Monoclinic, C_2
 $a = 20.847(2) \text{ \AA}$
 $b = 12.180(1) \text{ \AA}$
 $c = 7.1222(7) \text{ \AA}$
 $\beta = 101.738(2)^\circ$
 $V = 1770.5(3) \text{ \AA}^3$

$Z = 4$
 $D_x = 1.536 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation
 $\mu = 2.45 \text{ mm}^{-1}$
 $T = 295(2) \text{ K}$
 Block, yellow
 $0.26 \times 0.25 \times 0.24 \text{ mm}$

Data collection

Bruker SMART 1000 area-detector diffractometer
 φ and ω scans
 Absorption correction: none
 4679 measured reflections

3068 independent reflections
 2244 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.019$
 $\theta_{\text{max}} = 25.3^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.101$
 $S = 0.91$
 3068 reflections
 221 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0522P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.34 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983),
 with 1368 Friedel pairs
 Flack parameter: 0.05 (1)

H atoms were placed in calculated positions, with C—H 0.93–0.98 Å and $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{C})$, and were included in the refinement in the riding-model approximation.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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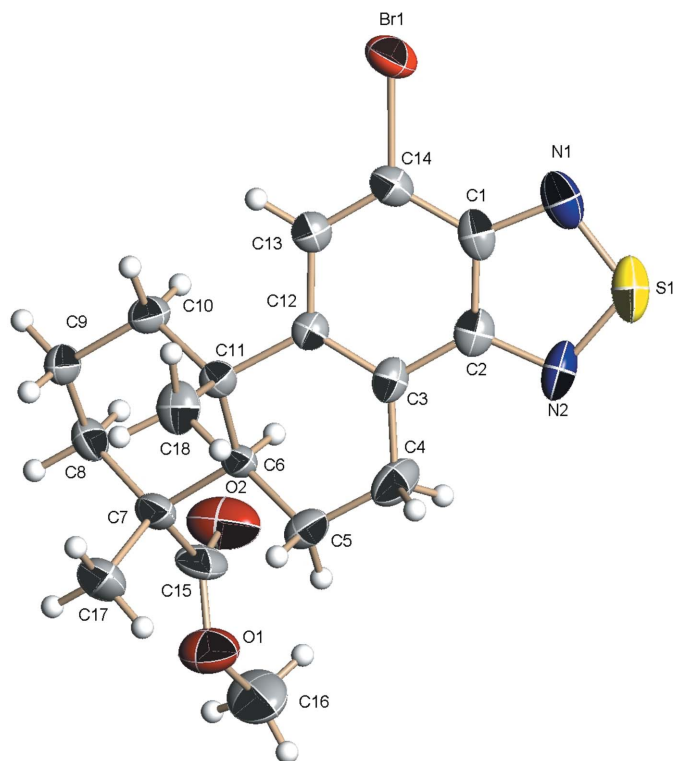


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as spheres of arbitrary radii.

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